



SEQUENCE LISTING

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Neidhardt, Helge

Paulista, Michael

<120> NEW GROWTH/DIFFERENTIATING FACTOR OF TGF- FAMILY

<130> 100564-09022

<140> US 09/386,450

<141> 1999-08-31

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<151> 1994-08-10

<150> DE P 43 26 829.3

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<160> 41

<170> PatentIn version 3.0

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<223> coding region for TGF-beta protein MP-52

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ggcaacagca gcgtgaagtt ggaggctggc ctggccaaca ccatcaccag cttattgac 1260

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<223> TGF-beta protein MP-52

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Leu Asp Leu Glu Phe Ile Cys Thr Val Leu Gly Ala Pro Asp Leu Gly

20 25 30

Gln Arg Pro Gln Gly Thr Arg Pro Gly Leu Ala Lys Ala Glu Ala Lys

35 40 45

Glu Arg Pro Pro Leu Ala Arg Asn Val Phe Arg Pro Gly Gly His Ser

50 55 60

Tyr Gly Gly Gly Ala Thr Asn Ala Asn Ala Arg Ala Lys Gly Gly Thr

65 70 75 80

Gly Gln Thr Gly Gly Leu Thr Gln Pro Lys Lys Asp Glu Pro Lys Lys

85 90 95

Leu Pro Pro Arg Pro Gly Gly Pro Glu Pro Lys Pro Gly His Pro Pro

100 105 110

Gln Thr Arg Gln Ala Thr Ala Arg Thr Val Thr Pro Lys Gly Gln Leu

115 120 125

Pro Gly Gly Lys Ala Pro Pro Lys Ala Gly Ser Val Pro Ser Ser Phe  
130 135 140

Leu Leu Lys Lys Ala Arg Glu Pro Gly Pro Pro Arg Glu Pro Lys Glu  
145 150 155 160

Pro Phe Arg Pro Pro Pro Ile Thr Pro His Glu Tyr Met Leu Ser Leu  
165 170 175

Tyr Arg Thr Leu Ser Asp Ala Asp Arg Lys Gly Gly Asn Ser Ser Val  
180 185 190

Lys Leu Glu Ala Gly Leu Ala Asn Thr Ile Thr Ser Phe Ile Asp Lys  
195 200 205

Gly Gln Asp Asp Arg Gly Pro Val Val Arg Lys Gln Arg Tyr Val Phe  
210 215 220

Asp Ile Ser Ala Leu Glu Lys Asp Gly Leu Leu Gly Ala Glu Leu Arg  
225 230 235 240

Ile Leu Arg Lys Lys Pro Ser Asp Thr Ala Lys Pro Ala Ala Pro Gly  
245 250 255

Gly Gly Arg Ala Ala Gln Leu Lys Leu Ser Ser Cys Pro Ser Gly Arg  
260 265 270

Gln Pro Ala Ser Leu Leu Asp Val Arg Ser Val Pro Gly Leu Asp Gly  
275 280 285

Ser Gly Trp Glu Val Phe Asp Ile Trp Lys Leu Phe Arg Asn Phe Lys  
290 295 300

Asn Ser Ala Gln Leu Cys Leu Glu Leu Glu Ala Trp Glu Arg Gly Arg  
305 310 315 320

Ala Val Asp Leu Arg Gly Leu Gly Phe Asp Arg Ala Ala Arg Gln Val

325 330 335

His Glu Lys Ala Leu Phe Leu Val Phe Gly Arg Thr Lys Lys Arg Asp

340 345 350

Leu Phe Phe Asn Glu Ile Lys Ala Arg Ser Gly Gln Asp Asp Lys Thr

355 360 365

Val Tyr Glu Tyr Leu Phe Ser Gln Arg Arg Lys Arg Arg Ala Pro Leu

370 375 380

Ala Thr Arg Gln Gly Lys Arg Pro Ser Lys Asn Leu Lys Ala Arg Cys

385 390 395 400

Ser Arg Lys Ala Leu His Val Asn Phe Lys Asp Met Gly Trp Asp Asp

405 410 415

Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu Gly Leu

420 425 430

Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His Ala Val

435 440 445

Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro Pro Thr

450 455 460

Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe Ile Asp

465 470 475 480

Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val Val Glu

485 490 495

Ser Cys Gly Cys Arg

500

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<211> 24

<212> DNA

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<222> (1)..(24)

<223> MP-52 adapter primer

<400> 3

agaattcgca tgccatgggc gacg

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<210> 4

<211> 23

<212> DNA

<213> Artificial

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<222> (1)..(23)

<223> MP-52 internal primer

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cttgagtacg aggctttcca ctg

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<210> 5

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attcgcatgc catggtcgac gaag

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<223> MP-52 internal primer

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ggagcccacg aatcatgcag tca

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<223> MP-52 internal primer

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acagcaggtg ggtggtgtgg act

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<223> adapter sequence

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44

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<212> DNA

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<222> (1)..(20)

<223> MP-52 internal primer

<400> 9

ccagcagccc atccttctcc

20

<210> 10

<211> 24

<212> DNA

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<223> MP-52 internal primer

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tccagggcac taatgtcaaa cacg

24

<210> 11

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<212> DNA

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actaatgtca aacacgtacc tctg

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<210> 12

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<212> DNA

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<222> (1)..(10)

<223> adapter

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<210> 13

<211> 102

<212> PRT

<213> Artificial

<220>

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<222> (1)..(102)

<223> MP-52

<400> 13

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1            5            10            15

Asp Trp Ile Ile Ala Pro Leu Glu Tyr Glu Ala Phe His Cys Glu Gly  
20            25            30

Leu Cys Glu Phe Pro Leu Arg Ser His Leu Glu Pro Thr Asn His Ala  
35            40            45

Val Ile Gln Thr Leu Met Asn Ser Met Asp Pro Glu Ser Thr Pro Pro  
50            55            60

Thr Cys Cys Val Pro Thr Arg Leu Ser Pro Ile Ser Ile Leu Phe Ile  
65            70            75            80



Asp Ser Ala Asn Asn Val Val Tyr Lys Gln Tyr Glu Asp Met Val Val  
85 90 95

Glu Ser Cys Gly Cys Arg  
100

<210> 14

<211> 101

<212> PRT

<213> Artificial

<220>

<221> DOMAIN

<222> (1)..(101)

<223> portion of BMP 2 corresponding to MP 52

<400> 14

Cys Lys Arg His Pro Leu Tyr Val Asp Phe Ser Asp Val Gly Trp Asn  
1 5 10 15

Asp Trp Ile Val Ala Pro Pro Gly Tyr His Ala Phe Tyr Cys His Gly  
20 25 30

Glu Cys Pro Phe Pro Leu Ala Asp His Leu Asn Ser Thr Asn His Ala

35

40

45

Ile Val Gln Thr Leu Val Asn Ser Val Asn Ser Lys Ile Pro Lys Ala

50

55

60

Cys Cys Val Pro Thr Glu Leu Ser Ala Ile Ser Met Leu Tyr Leu Asp

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80

Glu Asn Glu Lys Val Val Leu Lys Asn Tyr Gln Asp Met Val Val Glu

85

90

95

Gly Cys Gly Cys Arg

100

<210> 15

<211> 101

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<221> DOMAIN

<222> (1)..(101)

<223> portion of BMP 4 corresponding to MP 52

<400> 15

Cys Arg Arg His Ser Leu Tyr Val Asp Phe Ser Asp Val Gly Trp Asn

1 5 10 15

Asp Trp Ile Val Ala Pro Pro Gly Tyr Gln Ala Phe Tyr Cys His Gly

20 25 30

Asp Cys Pro Phe Pro Leu Ala Asp His Leu Asn Ser Thr Asn His Ala

35 40 45

Ile Val Gln Thr Leu Val Asn Ser Val Asn Ser Ser Ile Pro Lys Ala

50 55 60

Cys Cys Val Pro Thr Glu Leu Ser Ala Ile Ser Met Leu Tyr Leu Asp

65 70 75 80

Glu Tyr Asp Lys Val Val Leu Lys Asn Tyr Gln Glu Met Val Val Glu

85 90 95

Gly Cys Gly Cys Arg

100

<210> 16

<211> 102

<212> PRT

<213> Artificial

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<221> DOMAIN

<222> (1)..(102)

<223> portion of BMP 5 corresponding to MP 52

<400> 16

Cys Lys Lys His Glu Leu Tyr Val Ser Phe Arg Asp Leu Gly Trp Gln  
1            5            10            15

Asp Trp Ile Ile Ala Pro Glu Gly Tyr Ala Ala Phe Tyr Cys Asp Gly  
          20            25            30

Glu Cys Ser Phe Pro Leu Asn Ala His Met Asn Ala Thr Asn His Ala  
          35            40            45

Ile Val Gln Thr Leu Val His Leu Met Phe Pro Asp His Val Pro Lys  
          50            55            60

Pro Cys Cys Ala Pro Thr Lys Leu Asn Ala Ile Ser Val Leu Tyr Phe  
65            70            75            80

Asp Asp Ser Ser Asn Val Ile Leu Lys Lys Tyr Arg Asn Met Val Val  
          85            90            95

Arg Ser Cys Gly Cys His  
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<211> 102

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<222> (1)..(102)

<223> portion of BMP 6 corresponding to MP 52

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Cys Arg Lys His Glu Leu Tyr Val Ser Phe Gln Asp Leu Gly Trp Gln  
1 5 10 15

Asp Trp Ile Ile Ala Pro Lys Gly Tyr Ala Ala Asn Tyr Cys Asp Gly  
20 25 30

Glu Cys Ser Phe Pro Leu Asn Ala His Met Asn Ala Thr Asn His Ala  
35 40 45

Ile Val Gln Thr Leu Val His Leu Met Asn Pro Glu Tyr Val Pro Lys  
50 55 60

Pro Cys Cys Ala Pro Thr Lys Leu Asn Ala Ile Ser Val Leu Tyr Phe  
65 70 75 80

Asp Asp Asn Ser Asn Val Ile Leu Lys Lys Tyr Arg Asn Met Val Val  
85 90 95

Arg Ala Cys Gly Cys His

100

<210> 18

<211> 102

<212> PRT

<213> Artificial

<220>

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<222> (1)..(102)

<223> portion of BMP 7 corresponding to MP 52

<400> 18

Cys Lys Lys His Glu Leu Tyr Val Ser Phe Arg Asp Leu Gly Trp Gln

1            5            10            15

Asp Trp Ile Ile Ala Pro Glu Gly Tyr Ala Ala Tyr Tyr Cys Glu Gly

20            25            30

Glu Cys Ala Phe Pro Leu Asn Ser Tyr Met Asn Ala Thr Asn His Ala

35            40            45

Ile Val Gln Thr Leu Val His Phe Ile Asn Pro Glu Thr Val Pro Lys

50            55            60

Pro Cys Cys Ala Pro Thr Gln Leu Asn Ala Ile Ser Val Leu Tyr Phe  
65                70                75                80

Asp Asp Ser Ser Asn Val Ile Leu Lys Lys Tyr Arg Asn Met Val Val  
              85                90                95

Arg Ala Cys Gly Cys His  
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acgtgggggtg gaatgactgg at

22

<210> 21

<211> 22

<212> DNA

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<221> misc\_feature

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<223> portion of BMP 3 corresponding to primer OD

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atattggctg gagtgaatgg at

22

<210> 22

<211> 22

<212> DNA

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<222> (1)..(22)

<223> portion of BMP 4 corresponding to primer OD

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atgtgggctg gaatgactgg at

22

<210> 23

<211> 22

<212> DNA

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acctgggctg gcaggactgg at

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<210> 24

<211> 22

<212> DNA

<213> Artificial

<220>

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<222> (1)..(22)

<223> portion of TGF-beta-1 corresponding to primer OD

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aggacctcgg ctggaagtgg at

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<210> 25

<211> 22

<212> DNA

<213> Artificial

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<222> (1)..(22)

<223> portion of TGF-beta-2 corresponding to primer OD

<400> 25

gggatctagg gtggaaatgg at

22

<210> 26

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<223> portion of TGF-beta-3 corresponding to primer OD

<400> 26

aggatctggg ctggaagtgg gt

22

<210> 27

<211> 22

<212> DNA

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<222> (1)..(22)

<223> portion of Inhibin alpha corresponding to primer OD

<400> 27

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22

<210> 28

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<212> DNA

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<222> (1)..(22)

<223> portion of Inhibin beta-gamma corresponding to primer OD

<400> 28

acatcggctg gaatgactgg at

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<210> 29

<211> 22

<212> DNA

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<400> 29

tcacggctg gaacgactgg at

22

<210> 30

<211> 29

<212> DNA

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<223> Primer OID

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29

<210> 31

<211> 21

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<213> Artificial

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<222> (1)..(21)

<223> portion of BMP 2 corresponding to primer OID

<400> 31

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<210> 32

<211> 21

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<223> portion of BMP 3 corresponding to primer OID



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<210> 33

<211> 21

<212> DNA

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<222> (1)..(21)

<223> portion of BMP 4 corresponding to primer OID

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21

<210> 34

<211> 21

<212> DNA

<213> Artificial

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<222> (1)..(21)

<223> portion of BMP 7 corresponding to primer OID

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21

<210> 35

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<213> Artificial

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<222> (1)..(21)

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<222> (1)..(21)

<223> portion of TGF-beta-2 corresponding to primer OID

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21

<210> 37

<211> 21

<212> DNA

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<222> (1)..(21)

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21

<210> 38

<211> 21

<212> DNA

<213> Artificial

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<222> (1)..(21)

<223> portion of Inhibin alpha corresponding to primer OID

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21

<210> 39

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<222> (1)..(21)

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<223> Adapter

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